

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Mizzen, Lee

Anthony, Lawrence S.D. Wu, Huacheng Bill Siegel, Marvin

- (i1) TITLE OF INVENTION: IMMUNE RESPONSE USING COMPOSITIONS CONTAINING STRESS PROTEINS
- (iii) NUMBER OF SEQUENCES: 8
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: HAMILTON, BROOK, SMITH & REYNOLDS, P.C.
 - (B) STREET: TWO MILITIA DRIVE
 - (C) CITY: LEXINGTON
 - (D) STATE: MASSACHUSETTS
 - (E) COUNTRY: USA
 - (F) ZIP: 02173
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (V1) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/977,787
 - (B) FILING DATE: 25-NOV-1997
 - (C) CLASSIFICATION:
- (vii) PPIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/756,621
 - (B) FILING DATE: 26-NOV-1996
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Granahan, Patricia
 - (B) REGISTRATION NUMBER: 32,227
 - (C) REFERENCE/DOCKET NUMBER: STS96 02A
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (781) 861-6240
 - (B) TELEFAX: (781) 861-9540
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: peptide

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Cys Val Gln Ile Ala Ser Asn Glu Asn Met Glu Thr Met 1 $$ 10

- (2) INFORMATION FOR SEQ ID NO:3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: other nucleic acid
 - (A) DESCRIPTION: /desc = "DNA"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

AAAGAAGAAT TCAGGCGAAT C

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 - (A) DESCRIPTION: /desc = "DNA"

21

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GTTCCGATCA CTAGTCCCAC G	21
(2) INFORMATION FOR SEQ ID NO:5:	
 (i) SEQUENCE CHAPACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
CTGCTTGAAT TCAGCCAAGT G	21
(2) INFORMATION FOR SEQ ID NO:6:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
<pre>(1i) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
(mi) SEQUENCE DESCRIPTION: SEQ ID No:6:	
CTGTTGACTA GTGTTTCCTC C	21
(2) INFORMATION FOR SEQ ID NO:7:	
(A) LENGTH: 9 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: unknown (D) TOPOLOGY: unknown	
(1i) MOLECULE TYPE: peptide	
(mi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
Leu Pro Tyr Leu Gly Trp Leu Val Pro 1 5	

- (2) INFORMATION FOR SEQ ID NO:8:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 13 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Cys Lys Lys Leu Pro Tyr Leu Gly Trp Leu Val Pro 1 5